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TD6TTT-26288660

FIG. 2A CSG2

CAGGACTGCGTGTGCACGGACAAGGTGGACAACAACCCCTGCTCAACGTCAATCGCCTGC
 Q D C V C T D K V D N N T L L N V I A C

 ACCCAGTGCCCTGCAACACCTCCTGCAGCCCTGGGTTGGAACATCATGAGGCCCGGG
 T H V P C N T S C S P G F E L M E A P G

 GAGTGCTGTAAAGAGTGTGAACAGACGCACTGTATCATCAACGGCCGACAAACAGCAC
 E C C K K C E Q T H C I I K R P D N Q H

 GTCATCCTGAAGCCCGGACTTCAAGAGCGACCCGAAGAACAACATTCCTTCAGC
 V I L K P G D F K S D P K N N C T F F S

 TCGGTGAAGATCCACAACCAGCTCATCTCGTCCGTTTCCAAACATCACCTGCCCACTTT
 C V K I H N Q L I S S V S N I T C P N F

 GATGCCAGCATTTGCATCCCGGGCTCCATCACATTCATGCCCAATGGATGCTGCAAGACC
 D A S I C I P G S I T F M P N G C C K T

 TGCACCCCTCGCAATGAGACCAGGGTGCCCTGCTCCACCGTCCCGTCACCACGAGGTT
 C T P R N E T R V P C S T V P V T T E V

 TCGTACGCCCGGCTGCACCAAGACCGTCTCATGATCATGCTCCGGTCTCGGGGACA
 S Y A G C T K T V L M N H C S G S C G T

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FIG. 2B CSG2

TTTGTCACTCGGCCAAGGCCAGGCCCTGGACCACAGCTGCTCCTGCTGCAAAGAG
F V M Y S A K A Q A L D H S C S C C K E

GAGAAACCAGCCAGCGTGAGGTGCTCCTGAGCTGCCCAATGGCGGCTCGCTGACACAC
E K T S Q R E V V L S C P N G G S L T H

ACCTACACCCACATCGAGAGCTGCCAGTGCCAGGACACCGTCTGCGGGCTCCCCACCGGC
T Y T H I E S C Q C Q D T V C G L P T G

ACCTCCCCGGCCCGCGTTCCCTAGGCATCTGGGGAGCGGGTGAGCGGGTGGGCA
T S R R A R S P R H L G S G

CAGCCCCTTCACTGCCCCTCGACAGCTTTACCTCCCCCGACCCCTCTGAGCCTCCTAAGCT

CGGCTTCCTCTCTCAGATAATTATGTCTGAGTTTGTTCAGTCCTTGCTTTCCAATA

ATAAACTCAGGGGACATGCAAAAAATAAAAAA

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FIG. 3A CSG3

ATTGGTGCTACCTGGCTCTCCTGCTCTGCAGCTCTACAGGTAGGCCCCAGCAGAGGGAG
 TAGGGCTCGCCATGTTTCTGGTGAGCCCAATTGGGCTGATCTTGGGTGCTGAACAGCTAT
 TGGGTCCACCCAGTCCCTTTTCAGCTGCTGCTTAATGCCCTGCTCTCTCCCTGGCCCCACC
 TTATAGAGAGCCCCAAGAGCTCCTGTAAGAGGAGAACTCTATCTGTGTTTATAATCTT
 GCACGAGCACCAAGTCTCCCTGGGCTTTGTGAATGAACACTACATTTATCCCTTTCTCT
 GCCCCAACCAAACTCTTTCCCTTCAAGAGGGCCTGCCCTGGTTCCCTCCACCCAACTGC
 ACCATGAGATCGGTCCAAGAGTCCATTCCCCAGGTGGAGCCAACTGTCAGGGAGGTCTT
 TCCCACCAAAACATCTTTTCAGTTGCTGGGAGGTGACCATAGGGCTCTGCTTTTAAAGATAT
 GGCTGCTTCAAAGGCCAGAGTCACAGGAAGGACTTCTTCCAGGGAGATTAGTGGTGATGG
 AGAGGAGAGTTAAATGACCTCATGTCTCTTGTCTCCACGGTTTGTGAGTTTTCACCTC
 TTCTAATGCAAGGGTCTCACACTGTGAACCACTTAGGATGTGATCATCTTTCAGGTGGCCA
 GGAATGTGAATGCTCTTTGGCTCAGTTCATCTAAAAAAGATATCTATTTTGAAAGTTCTCA

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FIG. 3B CSG3

GAGTTGTACATATGTTTCACAGTACAGGATCTGTACATAAAAGTTTCTTTCCCTAAACCAT
TCACCAAGAGCCAATATCTAGGCATTTCCCTCGGTAGCACAAATTTTCTNATTGCTTAGAA
AATTGTCCCTCCCTGTTCTCTGTCTGNAGACTTAAGTGAGTTAGTCTTTTAAGGAAAGC
AACGCTCCCTCTGAAATGCTGTCTTTTTTCTGTGCGGAAATAGCTGCTCTTTTTCGGG
AGTTAGATGTATAGAGTGTTTGTATGTAAACATTTTCTTGTAGGCATCACCCATGAACANAG
ATATATTTCTATTTANTTANTATATGTGCACTTCAAGAAGTCACCTGTCAGAGAAATAAA
GAATTGTCTTAAATGTCATGATGAGATGTCTCTTTTGCAATTGCTTGGAAGGGGTACCT
AGAGCCAAGGAAATTGGCTCTGCTTTGGGAAAAATTTTGCTGTATTATAGTAAACATACA
AAGGATGTC

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FIG. 4 CSG4

ATGAGTCCTGTGAAAAACAATGTGGGCAGAGGCCCTAAACATCGCCCTGTGTAATGGAACC
M S P V K N N V G R G L N I A L V N G T

ACGGAGCTGTGCTGGACAGAGGCATTGTGACATGTACTCTGGAGATGTTATGCACCTA
T G A V L G Q K A F D M Y S G D V M H L

GTGAAATTCTTAAAGAAATTCCGGGGGTGCACTGGTGCTGGTGGCCTCCTACGACGAT
V K F L K E I P G G A L V L V A S Y D D

CCAGGACCAAAATGAACGATGAAGCAGGAACTCTTCTCTGACTTGGGAGTTCCTAC
P G T K M N D E S R K L F S D L G S S Y

GCAAACAACCTGGGCTTCCGGGACAGCTGGGTCTTCATAGGAGCCAAAGACCTCAGGGGT
A K Q L G F R D S W V F I G A K D L R G

AAAAGCCCCCTTTGAGCAGTTCTTAAAGAACAGCCCCAGACACAAACAATACGAGGATCG
K S P F E Q F L K N S P D T N K Y E G W

CCAGAGCTGCTGGAGATCGAGGGCTGCATGCCCCCGAAGCCATTTTAGGGTGGCTGTGGC
P E L L E M E G C M P P K P F

TCTTCTCAGCCAGGGCCTGAAGAAGYTCTCCTGCATTAGGAGTCANAGCCCCGCAG

GCTGNAGGAGGAGCAGGGGGTGCTGCGTGGAAGGTGCTGCAGGCCTTGCACGCTGTG

TCGGCCCT

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FIG. 5A CSG6

TGTCTACTAAGGTATTTACAACTTATGACACGAATGGTAGATACAGTGTAAAGTGCG
V Y S R Y F T T Y D T N G R Y S V K V R

GGCTCTGGAGGAGTTAACGCAGCCAGACGGAGAGTGATACCCAGCAGAGTGGAGCACT
A L G G V N A A R R R V I P Q Q S G A L

GTACATACCTGGCTGGATGAGAAATGATGAATACAAATCGAATCCACCAAGACCTGAAAT
Y I P G W I E N D E I Q W N P P R P E I

TAATAAGGATGATGTTCAACACAAGCAAGTGTTGTTTCAGCAGAAACATCCTCGGAGGCTC
N K D D V Q H K Q V C F S R T S S G G S

ATTTGTGGCTTCTGATGTCCCAAATGCTCCCATACCTGATCTCTTCCACCTGGCCAAAT
F V A S D V P N A P I P D L F P P G Q I

CACCGACCTGAAGCGGAAATTCACGGGGCAGTCTCATTAATCTGACTTGGACAGCTCC
T D L K A E I H G G S L I N L T W T A P

TGGGGATGATTATGACCATGGGAACAGCTCACAAAGTATATCATTCGAATAAGTACAAGTAT
G D D Y D H G T A H K Y I I R I S T S I

TCTTGATCTCAGAGACAAGTTCAATGAATCTCTTCAAGTGAATACTACTGCTCTCATCCC
L D L R D K F N E S L Q V N T T A L I P

FIG. 5B CSG6

AAAGGAAGCCAACTCTGAGGAAGTCTTTTGTGTTTAAACCAGAAACATTACTTTTGAAAA
 K E A N S E E V F L F K P E N I T F E N
 TGGCACAGATCTTTTCATGCTATTCAGGCTGTTGATAAGGTCGATCTGAAATCAGAAAT
 G T D L F I A I Q A V D K V D L K S E I
 ATCCAACATTGCACGAGTATCTTTGTTTATTCCTCCACAGACTCCGCCAGAGACACCTAG
 S N I A R V S L F I P P Q T P P E T P S
 TCCTGATGAAACGTCTGCTCCTGTCCTAATAATTATCATATCAACAGCACCATTCCTGGCA
 P D E T S A P C
 TTCACATTTTAAATTAATGTGGAAGTGGGTAGGAGAACTGCAGTGTCAATAGNCTAGG
 CGTGAATTTTGTGCGGTGAATAAATAATSATTCANCCCTTTTGTGRTTTATATAAAAAA
 CGGNTNCCCATTGGGNNNTNNGGGGGGNNNTTTTAA

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TABLE "2623660"

FIG. 6 CSG7

AGTCGCTCTCCTAGCCCTTCTCTGTGCTCACCCCTCTGGCAATGCCATTGAGCCAGGTC
V A L L A L L C A S P S G N A I Q A R S

TTCCTCCTATAGTGGAGAGTATGAGGTGGTGGTGGAAAGCGATTCTCTCATTTCTGGCAA
S S Y S G E Y G G G G K R F S H S G N

CCAGTGGACGGCCCCCATCACCGCCCTCCGGTCCGAGTCAACACATACTACATCGTAGG
Q L D G P I T A L R V R V N T Y Y I V G

TCTTCAGGTGCGCTATGGCAAGGTGTGGAGCGACTATGTGGGTGGTCGCAACGGAGACCT
L Q V R Y G K V W S D Y V G G R N G D L

GGAGGAGATCTTCTGCACCCCTGGGGAATCAGTGATCCAGGTTTCTGGGAAGTACAAGTG
E E I F L H P G E S V I Q V S G K Y K W

GTACCTGAAGAAGCTGGTATTGTGTGACAGACAAGGCCCGCTATCTGTCTTTTGGGAAGA
Y L K K L V F V T D K G R Y L S F G K D

CAGTGGCACAAGTTTCAATGCCCGTCCCTTGCACCCCAACACCGTGCTCCGCTTCATCAG
S G T S F N A V P L H P N T V L R F I S

TGGCCGCTCTGTTCTCTCATCGATGCCATTGGCCTGCACCTGGGATGTTTACCCACTAG
G R S G S L I D A I G L H W D V Y P T S
CTGCAGCAGATGCTGAGCCTCCTCTCTTGGCAGGGGCACTGTGATGAGGAGTAAGAACT
C S R

CCTTATCACTAACCCCATC

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FIG. 7

FIG. 7 CSG8

TAAACTGCTGTTTGTTCCCTGTCCTGTCCTTTGGTTGGTATTTCAGTAAGTTTTCGGT
ATTCTCAAATTTTATCTAAATGGATAAACTATTAAACATAGAACATAAACCCCAATTCTCC
ATTTCATTTTCTCTTAGGCAATGAATCATACAAAACCTCAATATAGAGCAATGTTGTAAAT
GAATTGTTCTATTAAACAAAGAGGAGGTTCTAAGATATAAAGCCTCAGAGAACAGGAAGAA
AAGCGGGTCCATAAGAAGATGAGGTCTAACCGGGAAGATGCTGCTGAGAAGGCAGAGAC
AGATGTGGAAGAAATCTATCACCCAGTCATGTGCACCTGAATGTTCCACTGAAGTGCCAGT
TTACGACAAGGATGAAGTCTTTTCATTTTTCATGTTTATAGCAAGCCATTCTCTAAACAGC
CCAACTGGCATTTAAATTACCCAATACTGTATATAAGGCCAAATATGGACAGTTACTTTCCT
CTTGCCCTGTTTCATATCCTTCAGTGACATTGAGGAAGCAGTGTTCCTCTTTTAAAGGGGA
ATAGTGTCAACCTTTCATCTCTTACATCTTTACCCCTCTCTCTTTTTCCTTTG
ATTTTCCCCCTTATTGATGGGACTGATATTTCATCTCTGTTTTCATGAAACATTTGGAAACT
GTCCGGGCTTTTATTAAAGCTCTGTAGAAATTAAATGTTCTGGAATTAT

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FIG. 8 CSG9

CAGGAGGAGAGCCCTTCCCCAAGCAACAATCCAGAGCAGCTGTGCAACAACGGTGCCAT

AAATAAGGCCCTCCTGGACCATGAATGCGAGTCCGCTGAGCTGCGTACCGGAGCCCCACGGT

GGTCATGGCTGCCAGAGCGCTCTGCATGCTGGGGCTGGTCTGCGCCTTGCTGTCTCCTCCAG
M A A R A L C M L G L V L A L L S S S

CTCTGCTGAGGAGTACGTGGGCCCTGTCTGCAAAACCAGTGTGCCCGTGCCAGCCAAGGACAG
S A E E Y V G L S A N Q C A V P A K D R

GGTGACTGCGGCTACCCCCCATGTCACCCCCCAAGGAGTGCAACAACCGGGCTGCTGCTT
V D C G Y P H V T P K E C N N R G C C F

TGACTCCAGGATCCCTGGAGTGCCCTTGGTGTTTCAAGCCCCCTGACAGGGAAGCAGGAATG
D S R I P G V P W C F K P L T G K Q E C

CACCTTCTGAGGCACCTCCAGCTGCCCCCCCGCGGGGATGCGAGGCTCGGAGCACCCCT
T F *

TGCCCGGCTGTGATTGCTGCCAGGCACTGTTCATCTCAGCTTTTCTGTCCCTTTGCTCCC

GGAAGCGCTTCTGCTGAAAGTTTCATATCTGGAGCCTGATGTTTAACTAGTCCCCATGCTC

CACCCGAAAAAATAAAAAAAAAAAAAA

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FIG. 9A CSG10

AAGCTCTTCTACAGGACCAGCCACTAGCGCAGCTCGAGCGATGGCCTATGTCCCCGCAC
M A Y V P A P

CGGGCTACCAGCCACCTACAACCCGACGCTGCCCTTACTACCAGCCCATCCCGGGCGGGC
G Y Q P T Y N P T L P Y Y Q P I P G G L

TCAACGTGGGAATGTCTGTATTACATCCAAGAGTGGCCAGCGAGCACATGAAGCGGTTCT
N V G M S V Y I Q G V A S E H M K R F F

TCGTGAACCTTGTGTTGGCAGGATCCGGGCTCAGACGTGCGCTTCCACTTCAATCCGC
V N F V V G Q D P G S D V A F H F N P R

GGTTGACGGCTGGGACAAGGTGCTTCAACACGTTCAGGGCGGGAAGTGGGCGACGC
F D G W D K V V F N T L Q G G K W G S E

AGGAGAGGAAGAGGAGCATGCCCTTCAAAAGGGTCCCGCCTTTGAGCTGGTCTTCATAG
E R K R S M P F K K G A A F E L V F I V

TCCTGGCTGAGCACTACAAGGTGGTAAATGGAATCCCTTCTATGATACGGGCACC
L A E H Y K V V N G N P F Y E Y G H R
GGCTTCCCCTACAGATGATCACCCACCTGCAAGTGGATGGGATCTGCAACTTCAATCAA
L P L Q M V T H L Q V D G D L Q L Q S I

TCAACTTCATCGGAGGCCAGCCCTCCGGCCCCAGGACCCCGATGATGCCACCTTACC
N F I G G Q P L R P Q G P P M M P P Y P

FIG. 9B

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FIG. 9B CSG10

CTGGTCCCGGACATTGCCCATCAACAGCTGAACAGCCTGCCCCACCATGGAAGACCCCAA
G P G H C H Q Q L N S L P T M E G P P T
CCTTCAACCCCGCCTGTGCCATATTTCCGGAGGCTGCAAGAGGCTCACAGCTCGAAGAA
F N P P V P Y F G R L Q G G L T A R R T
CCATCATCAAGGGCTATGTGCCCTCCACAGGCAAGAGCTTTGCTATCAACTTCAAGG
I I I K G Y V P P T G K S F A I N F K V
TGGCTCCTCAGGGACATAGCTCTGCCACATTAATCCCCGCATGGCAACGGTACCGTGG
G S S G D I A L H I N P R M G N G T V V
TCCGGAACAGCCTTCTGAATGGCTCGTGGGATCCGAGGAGAAGATCACCCACAACC
R N S L L N G S W G S E E K K I T H N P
CATTGGTCCCGGACAGTTCTTTGATCTGTCCATTCCGCTGTGCTTGATCGCTTCAAGG
F G P G Q F F D L S I R C G L D R F K V
TTTACGCCAATGGCCAGCACCTCTTTGACTTTTGCCCATCGCCCTCTCGGCCTTCCAGAGGG
Y A N G Q H L F D F A H R L S A F Q R V
TGGACACATGGAAATCCAGGGTGATGTCACCTTGTCCTATGTCCAGATCTAATCTATTC
D T L E I Q G D V T L S Y V Q I
CTGGGCCATAACTCATGGGAAAACAGAAATTATCCCCCTAGGACTCCTTTCTAAGCCCCCTA
ATAAAATGTCTGAGGGTGTCTCAAAAAAATAAAAAA

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TABLE "E" 223355D

FIG. 10 CSG11

GTTGATATTAAACCAAGTGAACCAACATGACACCTCTCTGAAACCTATTAGTGTCTCC
V D I K T S E T K H D T S L K P I S V S

TACAACCCAGCCACAGCCAAAGAAATTATCAATGTGGGCATTCTTCCATGTAAATTTT
Y N P A T A K E I I N V G H S F H V N F

GAGGACAACGATAACCGATCAGTGCTGAAGGTGTCCTTTCTCTGACAGCTACAGGCTC
E D N D N R S V L K G G P F S D S Y R L

TTTCAGTTCATTTCCTGCTGGGCGAGTACAAATGAGCATGGTTCAGAACATACAGTGGAT
F Q F H F H W G S T N E H G S E H T V D

GGAGTCAAAATATTCTGCCGAGCTTCACGTGGCTCACTGGAATTCTGCAAGTACTCCAGC
G V K Y S A E L H V A H W N S A K Y S S
CTTGCTGAAGCTGCCCTCAAAGGCTGATGGTTGGCAGTTATTGGTGTTTTGATGAAGGT
L A E A A S K A D G L A V I G V L M K V
GGTGAGGCCCAACCCAAAGCTGCAGAAAGTACTTGATGCCCTCCCAAGCAATTAAACCAAG
G E A N P K L Q K V L D A L Q A I K T K
GGCAAACGAGCCCCATTCAACAAATTTTGACCCCTCTACTCTCCTTCCTTCATCCCTGGAT
G K R A P F T N F D P S T L L P S S L D

TTCTGGACCTACCCCTGGCTCTCTGACTCATCCTCCTCTTTATGAGAGTGTAACTTGGATC
F W T Y P G S L T H P P L Y E S V T W I
ATCTGTAAAGGAGAGCATCAGTGTCAAGTTCAGAGCAGTTGGCACAATTCCGGAGCCTTCTA
I C K E S I S V S S E Q L A Q F R S L L

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FIG. 11 CSG12

CGCTCCGGCGGGCGTGGCCAGTACTAGAAAGCGGAGCGCGCGGACCATGGCGCG
G S G R A W P V T R R R G A A G P W R R

GCGGCGAGCGGAGTCCAGAGCGGAGAACGAGGAGGAGGAGCAGTTGCTTCT
R R T S G V Q R R E D E E E E Q L V L

GGTGGAATTATCAGGAATTATTGATTCAGACTTCCTCTCAAAATGTGAAAATAATGCAA
V E L S G I I D S D F L S K C E N K C K

GGTTTGGGCATTGACACTGAGAGGCCCATTCCTGCAATGGACAGCTGTGCTTTGCTGG
V L G I D T E R P I L A M D S C V F A G

GGAGTATGAGACACTCTAGGGACCTGTGTTATATTGAAGAAAATGTTGACATGCTGA
E Y E D T L G T C V I F E E N V E H A D

TACAGAAGGCAATAATAAACAGTGTCTAAATATAATGCCCATACAAATGAAGAAGCTCAG
T E G N N K T V L K Y K C H T M K K L S

CATGACAAGAACTCTCCTGACAGAGAAGGAGAGAAGAAACATAGTGGGTGGA
M T R T L L T E K K E G E E N I G G V E

ATGGCTGCAATAAGGATATGGTTTCTCCCTTTGACCCCAACAGGTTTGTAACTTTTCTA
W L Q I R I W F L P L T Q Q V C

CCATGAAATTGAGGACGAGGAAGTGGTAGCTTTCAGCCCCCGTTAAATCTTTGGATTTCGG
AGGGGGTGGGTTTCAATG

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FIG. 12 CSG13

GTGGCAGAAAGATAGGTTGGAGACAATTGATTCCTCGATGATATAAAAATGTTAAGTA

CCATGAATGNAATGCTGTTAGGCTGGAATGCGCCAAGATAAAAGGTGGGCGCATGGCATCAA

AAGGTAGGTCAACATATTAAATAATTCCCATGTATTGAAATATCCAGAAATATATAGACA

GATCTATAGAGATAGAAACTGGTCTGCCCCAGGACTAGGGGTGTCTAAGGATAAGGAGCT

TCTTTTGGATGGTGAAATAACCTAAAAATATATTGTGCCCATTTGTTGCCACAACCTTTGTG

GAATATATTAAACCGGTTAATTGTACTCACTAAATGTCCCTCTCTTAAATTTAAGC

TGTTTNCCTGGACAGAAAGGGAAGNNACCAAGGGGNAAAAAATTTT

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FIG. 13 CSG14

GCCCTGGGCTTTGGGGGCTCCAAACATGGTATGCAGAAATGTGATGTTACAGGTCAG
TACAAACCTCAGTCCTTAGAAACCCCTCCACACTTCAGCTCTGCCACCCACTTTCTCTGTCATT
TATTTATATAGGACTGTAGTTTTTTTTTAGTTTCGAGAGCCCTTTCGAAGCTTAATTTATATT
CTTCTTTGTACCTTTTTTCTAAATTACCAAGATATTACACAAAGGTAAATTAAATGTT
CTCTGTTTTATGCTTTATCTGATGGAGGCAAAATATCCTCTTATTGTTGATCAAAGGGGC
AAAAGAAATTTAGAGGCAAAATGAACAAGCGATAGGCTATTGCCAACCTGAGAAAGAGAACTG
NTCCTTCCATCGTAAATTTAGNAGNCCAAAGTAGGTAATGGGAACCAAGTTGTTACTTTT
TTCTAGTAGTTATTTTCCCTTTTTTNNTTTTTTGTGGTACCCTCTTACAGNGNCCCAAACCT
CCATTCTCTTTAAAGGGGTTTTTATGGGGGCTTACTGCAGGTTAAAAATTGGGGNCCAC
CATTTTTAAAGGGGCTACCAGAGGGAGGGGGTCCCCNTTNCNAAAAAAATG